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RAW SEQUENCE LISTING DATE: 07/12/2000
PATENT APPLICATION: US/09/606,977 TIME: 11:29:16Input Set : D:\Maizestcs.rpt
Output Set: N:\CRF3\07122000\I606977.raw

1 <110> APPLICANT: Byrum, Joseph R.
3 <120> TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
4 Plants
6 <130> FILE REFERENCE: 38-21(15877)B
C--> 8 <140> CURRENT APPLICATION NUMBER: US/09/606,977
C--> 8 <141> CURRENT FILING DATE: 2000-06-28
8 <160> NUMBER OF SEQ ID NOS: 82359
10 <210> SEQ ID NO: 1
11 <211> LENGTH: 280
12 <212> TYPE: DNA
13 <213> ORGANISM: Zea mays
15 <223> OTHER INFORMATION: unsure at all n locations
W--> 16 <400> SEQUENCE: 1
18 ctctggtg gagggttc atacccgtcg gagtaaaaacc ggagttctgg ttgttaatggc 60
20 accctgtgta cgagtgtctg ttcatgttgg ttgtcgaacgc ctgaaacattt gggtatgg 120
W--> 22 agtgaatac gngtttcttc cttatattcga gcactaagac tcgcctgtcg gctagctgaa 180
24 ccgcctaacc gagcgtgatg tgccctgcgc ggaagggtgac gatgtgaggtt tctgtattcc 240
26 ggaggcgtaa gagtttccctcg gtcgggtcg gtttgcctt 280
29 <210> SEQ ID NO: 2
30 <211> LENGTH: 360
31 <212> TYPE: DNA
32 <213> ORGANISM: Zea mays
34 <400> SEQUENCE: 2
36 agagagcgtg ccaaaaaaaa gttgcgggtt gatcttattt atacgcctag tgcgcgtaaaa 60
38 attgaagac cccgcgtcgtc agtgaatgtt gtattcttag caaaggaaag gtgtttttc 120
40 ggacccttcgg cgtaaagac tcgtccatgt cgcacatctaa atttattttt ttaaaaataaa 180
42 ttaaatatgc gaggggctac tggggggc cttctgttta caaagggtcct caaaaacagg 240
44 atttaacaat atttctggag tataatgtgt gacgaggat ttcggactc aagtcgttat 300
46 cacaacagga gaagaccaga atagtagcga cgttgatatac gcgcggccaaa tgtgaacagg 360
49 <210> SEQ ID NO: 3
50 <211> LENGTH: 433
51 <212> TYPE: DNA
52 <213> ORGANISM: Zea mays
54 <223> OTHER INFORMATION: unsure at all n locations
W--> 55 <400> SEQUENCE: 3
W--> 57 agcttnggag ccgaagcaac tccagctttg tacgatttca tcagctcaga aagtggaaat 60
59 gagttcacccg gatgtggcgc ccaatgttgg ggacttggc tcaaatgctt cgaattaaga 120
61 acaagacaac acaaaaaatgt taagtataaa ttcccttcgt ccgtcgtaaacgattatctccc 180
63 caaggattta atgacttcgt gacgaaaggcc ataaataaaatgatc tttcacgaaatgttcc 240
65 cgtgatcata ttttaaagac aacaatggaa aatgaaatat aagcacaacaa atatattc 300
67 atattctgtt atatgaaatat acaagtattt aacatataaca ttatcgaaca ttatatacc 360
W--> 69 catcttgca gaaaacaata atccggcgt aacgtgtncg tgatatacaga tcgcgtgaac 420
71 agtaccggga tac 433
74 <210> SEQ ID NO: 4
75 <211> LENGTH: 359
76 <212> TYPE: DNA
77 <213> ORGANISM: Zea mays

ENTERED PS

RAW SEQUENCE LISTING

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Input Set : D:\Maizestcs.rpt

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79 <400> SEQUENCE: 4
81 aactagctaa ttgatataga tccttggta gtcgcacgtt tgactttgtt gatcccaacc 60
83 caaacatcgc ttttaaggaa tattgcctta ttaatggaat actgactaga aacctatgtt 120
85 gccagggtta cccggaaa gtttattaag aataacttac tatcttctat ataactgata 180
87 ttatagcata catgagccatc atgttatgtt tggctgtttaa aaacaagaaa agaattgtca 240
89 aaaaacgact gaccaccgccc agactcacct gcggaaatgtt tattttttt gggacaatcc 300
91 aatacttacaa ggcataccctt ctgggtttaa acttataactt gtgagcttag gaagagtgg 359
94 <210> SEQ ID NO: 5
95 <211> LENGTH: 358
96 <212> TYPE: DNA
97 <213> ORGANISM: Zea mays
99 <223> OTHER INFORMATION: unsure at all n locations
W--> 100 <400> SEQUENCE: 5
102 gtttattgtt gaagcaccag aagccaatc aaggaaggaa tccatgtggg agttcgcaag 60
104 aggaaaagttt gcgtgtcgcc atcaatgtt ccataataag ctatgtgc aagcggaaacct 120
W--> 106 agaaggcttgc agatgattn ttgagttagcc agaatcagcg ttggcaatca gcaacaggat 180
108 gctccctaa gaccaggat ttgttgaatgtt ccatctctc gaagaggatgc aaagtgaaga 240
110 tatgtatgtt aagttaaatgtt ataccatataa cccttctaa acgaatctgc aggacgacat 300
112 ttctttaatgtt gggtaggatt ttagcatcccc acaaataaaaa ttctgaatattt tctcaact 358
115 <210> SEQ ID NO: 6
116 <211> LENGTH: 449
117 <212> TYPE: DNA
118 <213> ORGANISM: Zea mays
120 <223> OTHER INFORMATION: unsure at all n locations
W--> 121 <400> SEQUENCE: 6
123 taaggggaaa cactgatgtt tggaaagga aatgtgcctt ttggccattt ctatgtat 60
W--> 125 tggatgggg gtcacac aatgtcttaa gtgtgaatnt atgtctatgg ataaacaaaag 120
127 cgcaatcaaa gagaaaaggt atgtttctaa gtcctagttc attgggttttg tgactaata 180
129 tacttgtctaa agttttggaa acagaaatgtt taagaaaaga gaagaggatgc aaaaatgtcg 240
131 gtctggagca cccggactgtc cgggtgtca cccggacgtt tccgggtgc caggctgtact 300
133 cggcgcgaac tcggccgtt caggatatgc cccggcgtt acggctaaaa ttacccggac 360
135 tattccgtgtt gacccggact gtccgggtt ctaacggatgc actgagccaa cggtccggcag 420
137 cggaatctgc ggcgcacac tggctggcc 449
140 <210> SEQ ID NO: 7
141 <211> LENGTH: 344
142 <212> TYPE: DNA
143 <213> ORGANISM: Zea mays
145 <400> SEQUENCE: 7
147 ttatgttata gtttactatc ttatgggatt ttttcaaaca agtcaaccat gcccgtggaaa 60
149 ttgttctca agttggcat aattgccttt tagatgtac taccttggaaa agacttgc 120
151 ttatgttataa aaatatgtaa attttgcattt cggggatgtt gtaatgtt aatgtctt 180
153 taaaactgcgatgtt acaggatcaa acctaataa aatgtctt aacagcgatc 240
155 ggatgtatgtt tagatgtt cccactaatt agacatgtt aggtacattt gtaaaaaat 300
157 atatgttataa atatgttataa aatgtctt ccaatataatg aaaa 344
160 <210> SEQ ID NO: 8
161 <211> LENGTH: 186
162 <212> TYPE: DNA
163 <213> ORGANISM: Zea mays
165 <400> SEQUENCE: 8

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```

167 ccagtaggct actataacag tgatatcact ttatcaatta cagatagtga cctcacactc 60
169 ttgaggatg attcaacttt taagaagaac actatatacta aagaaaggta cccattttaa 120
171 taatgcacc gagcagttaa ttaagattac aaaaaattga ggagggggga aatacaattt 180
173 ccacct 186
176 <210> SEQ ID NO: 9
177 <211> LENGTH: 404
178 <212> TYPE: DNA
179 <213> ORGANISM: Zea mays
181 <223> OTHER INFORMATION: unsure at all n locations
W--> 182 <400> SEQUENCE: 9
W--> 184 tgggtcccta agggaaattgt tactaacctt gnaggatcca acaagagttt ggtacctaag 60
186 acccaacgtt aaatttgcgt tgcagggtta tgcattccggg ggctcaagggtt ggattatccga 120
188 cagcgatgc acaaaccata tgacggggga gaagaagatg ttacacttc acgtcaagaa 180
190 taaggattcc caagatcaa tcatattcgg ggttggaaac caaaggagg ttgaaagggtt 240
192 gggcaaggatt gcaatataca atgagactc tatctctaat gtgttttag ttgagtctct 300
194 tggatataat ttactatctg tcagtcaattt atgttaatgtt ggttattttt tgctatttac 360
196 aaatgttaat gtgtctgtct ttataagaag tgatggttca ctat 404
199 <210> SEQ ID NO: 10
200 <211> LENGTH: 343
201 <212> TYPE: DNA
202 <213> ORGANISM: Zea mays
204 <223> OTHER INFORMATION: unsure at all n locations
W--> 205 <400> SEQUENCE: 10
207 ggagccgacg cattgcctca cggagaagc gtatttactc tcgtcgact tcggcttcc 60
209 gcccggcac gtggaccggc ctaaccagggt agtcttcgtc caattccctt cacccacaac 120
211 ttccctgaca tccccgtaaag ctcttcacc cgttcaattt aactatcgcc cggtgatgtt 180
213 gccggactcc tcggccggca cgagcatctc cgcctgcga cgtggaccga ccgactccaa 240
W--> 215 ccatccccga cggcggcccg cacatcggtt tgatccctca gaccttccct acgtntctca 300
217 ccacccctacc ggagaaatctt cggccgggtt aagccctcc gcc 343
220 <210> SEQ ID NO: 11
221 <211> LENGTH: 392
222 <212> TYPE: DNA
223 <213> ORGANISM: Zea mays
225 <223> OTHER INFORMATION: unsure at all n locations
W--> 226 <400> SEQUENCE: 11
W--> 228 tgcaagcttc ggctatgtct aannaatttg cagttaaaca atgcaatagc aatgaatgtc 60
W--> 230 gtggtaactt cacacccatcc cgtctgtttt tatagtgtcc cangtggaa ggttaatccg 120
232 caaggcttcc agcaccgaac aaacagtcac ccgcactcac tgagcggtt gcccacagggt 180
234 tcgagaaggtaa gatcgctgg cgcgcgcgtt attgggtccaa gatggggcgc gcgcgcacat 240
236 tattttatcc gtttctcgcc aacgagctca gggaaagggtt tttttggatc ttccggctcc 300
238 cgaagcttag aagactttt caccgttacg gctcgatcc aaaaacgttcc tatcaccgcg 360
240 aatgggttac ttgtgggggtt attgttttgtt ac 392
243 <210> SEQ ID NO: 12
244 <211> LENGTH: 447
245 <212> TYPE: DNA
246 <213> ORGANISM: Zea mays
248 <223> OTHER INFORMATION: unsure at all n locations
W--> 249 <400> SEQUENCE: 12
W--> 251 tctaaaggga gaacacgtga taggtttgac taacgtgttcat ttcaanaag atagacccgtt 60

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343 cacggatctg gttcaagaat tcttagcatt caaatgttcc cctactctga aaaaatggga 180
345 aatgc当地 aataaggggg agaagaagga aggggatctt gtccggcttc cttaccactt 240
347 caaatttc当地 aataattttta agaaagccctg ccaagagtgg ctggatcacag ttgaggttgat 300
349 gtgtatgaa attcttggaa attttcgaa aaaagaagat caactgtatga cagcagcatt 360
351 ctgcaccctg cccaagc当地 ggctgaaccg agtgctgtat gcactgggtt tcgaatatcc 420
353 tgattat 427
356 <210> SEQ ID NO: 17
357 <211> LENGTH: 211
358 <212> TYPE: DNA
359 <213> ORGANISM: Zea mays
361 <223> OTHER INFORMATION: unsure at all n locations
W--> 362 <400> SEQUENCE: 17
W--> 364 ntgc当地 cacaacat tataacttca tgaactttt atactgtca ggtactacgg 60
366 tcttagggc当地 tccatgtat ttc当地 agattt ctgtgatcaa taaaatggcg atgtgttgaa 120
368 catcctc当地 cttgttgc当地 ctaataactt gaaccattt cggaatctc tcatccaca 180
370 caaaaatgtc atgtcttctt ttttggagacc t 211
373 <210> SEQ ID NO: 18
374 <211> LENGTH: 390
375 <212> TYPE: DNA
376 <213> ORGANISM: Zea mays
378 <223> OTHER INFORMATION: unsure at all n locations
W--> 379 <400> SEQUENCE: 18
381 agctggc当地 tccgc当地 gtgc cccctgctt actgc当地 tctcataactc tccctgegt 60
383 cctgctcggt tccctgctc gccc当地 cc当地 ggaa aacttcccttcc tgc当地 gtc当地 gccc当地 cc当地 120
385 cc当地 cgttcc tccatgtatc tgc当地 gagatcc gccc当地 cc当地 180
387 acgc当地 caagaaat ttttccctgc gtc当地 ggcc当地 gagatccat tccctctgc 240
W--> 389 acgaatncat gtc当地 ccccgccg cccccc当地 gagatccatc tgc当地 gtgttctgc 300
391 cagcgtc当地 tgc当地 gagatccatc tgc当地 ctgc当地 cccatc当地 cgc当地 ccatc当地 gatcc 360
393 ccttgc当地 gggttccatgc当地 cccctggctcg 390
396 <210> SEQ ID NO: 19
397 <211> LENGTH: 373
398 <212> TYPE: DNA
399 <213> ORGANISM: Zea mays
401 <400> SEQUENCE: 19
403 cttc当地 cccat tccatgc当地 ctgc当地 gtc当地 catgaaatgc aatacgc当地 ctgc当地 tgatccgaa 60
405 gtatatttcat gcatagatat gtc当地 gatggatgaa gaatttgc当地 ttatgc当地 gtc当地 ttatgc当地 120
407 accaactgact gcatatgttgg ggtatatttcat gtc当地 aacaata tcaaaacacaa cctgttaact 180
409 tctt当地 tatttgc当地 gacatgc当地 tgc当地 gagatccatc tgc当地 ttatgc当地 gagatccat 240
411 ctgc当地 ctttccatc cccaaaccac aaagagggttgc tgc当地 atttgc当地 gagatccat 300
413 ttgc当地 atttgc当地 tccatgc当地 cccatc当地 cgc当地 ccatc当地 gatcc 360
415 caatgtggac tat 373
418 <210> SEQ ID NO: 20
419 <211> LENGTH: 400
420 <212> TYPE: DNA
421 <213> ORGANISM: Zea mays
423 <223> OTHER INFORMATION: unsure at all n locations
W--> 424 <400> SEQUENCE: 20
426 cacaacttatttgc ctatgtc当地 ttttgc当地 ggtctttaat gctaccctac atggcatgtt 60
428 tccttactca tatgggttgc当地 taggc当地 accgatc当地 tctattaaga attctattaa agatatgagc 120

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JULY 1

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

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Input Set : D:\Maizestcs.rpt
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L:8 M:270 C: Current Application Number differs, Replaced Current Application No
L:8 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:16 M:283 W: Missing Blank Line separator, <400> field identifier
L:22 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:1
L:22 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:1
L:22 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:1
L:22 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:1
L:55 M:283 W: Missing Blank Line separator, <400> field identifier
L:57 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:3
L:57 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:3
L:57 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:3
L:57 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:3
L:69 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:3
L:69 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:3
L:69 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:3
M:340 Repeated in SeqNo=3
L:100 M:283 W: Missing Blank Line separator, <400> field identifier
L:106 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:5
L:106 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:5
L:106 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:5
L:106 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:5
L:121 M:283 W: Missing Blank Line separator, <400> field identifier
L:125 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:6
L:125 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:6
L:125 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:6
L:125 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:6
L:182 M:283 W: Missing Blank Line separator, <400> field identifier
L:184 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:9
L:184 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:9
L:184 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:9
L:184 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:9
L:205 M:283 W: Missing Blank Line separator, <400> field identifier
L:215 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:10
L:215 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:10
L:215 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:10
L:215 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:10
L:226 M:283 W: Missing Blank Line separator, <400> field identifier
L:228 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:11
L:228 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:11
L:228 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:11
L:228 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:11
L:230 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:11
L:230 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:11
L:230 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:11
M:340 Repeated in SeqNo=11
L:249 M:283 W: Missing Blank Line separator, <400> field identifier
L:251 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:12
L:251 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:12

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L:251 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:12
L:251 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:12
L:362 M:283 W: Missing Blank Line separator, <400> field identifier
L:364 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:17
L:364 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:17
L:364 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:17
L:364 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:17
L:379 M:283 W: Missing Blank Line separator, <400> field identifier
L:389 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:18
L:389 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:18
L:389 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:18
L:389 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:18
L:424 M:283 W: Missing Blank Line separator, <400> field identifier
L:436 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:20
L:436 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:20
L:436 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:20
L:436 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:20
L:447 M:283 W: Missing Blank Line separator, <400> field identifier
L:461 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:21
L:461 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:21
L:461 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:21
L:461 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:21
L:472 M:283 W: Missing Blank Line separator, <400> field identifier
L:478 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:22
L:478 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:22
L:478 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:22
L:478 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:22
L:517 M:283 W: Missing Blank Line separator, <400> field identifier
L:527 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:24
L:527 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:24
L:527 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:24
L:527 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:24
L:540 M:283 W: Missing Blank Line separator, <400> field identifier
L:554 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:25
L:554 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:25
L:554 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:25
L:565 M:283 W: Missing Blank Line separator, <400> field identifier
L:567 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:26
M:340 Repeated in SeqNo=26
L:606 M:283 W: Missing Blank Line separator, <400> field identifier
L:614 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:28
L:625 M:283 W: Missing Blank Line separator, <400> field identifier
L:627 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:29
L:670 M:283 W: Missing Blank Line separator, <400> field identifier
L:672 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:31
M:340 Repeated in SeqNo=31
L:693 M:283 W: Missing Blank Line separator, <400> field identifier
L:695 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:32
M:340 Repeated in SeqNo=32

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L:716 M:283 W: Missing Blank Line separator, <400> field identifier
L:718 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:33
L:741 M:283 W: Missing Blank Line separator, <400> field identifier
L:753 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:34
L:764 M:283 W: Missing Blank Line separator, <400> field identifier
L:766 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:35
L:861 M:283 W: Missing Blank Line separator, <400> field identifier
L:875 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:40
L:908 M:283 W: Missing Blank Line separator, <400> field identifier
L:918 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:42
L:933 M:283 W: Missing Blank Line separator, <400> field identifier
L:937 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:43
M:340 Repeated in SeqNo=43
L:1014 M:283 W: Missing Blank Line separator, <400> field identifier
L:1018 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:47
L:1033 M:283 W: Missing Blank Line separator, <400> field identifier
L:1045 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:48
L:1054 M:283 W: Missing Blank Line separator, <400> field identifier
L:1058 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:49
M:340 Repeated in SeqNo=49
L:1099 M:283 W: Missing Blank Line separator, <400> field identifier
L:1107 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:51
M:340 Repeated in SeqNo=51
L:1124 M:283 W: Missing Blank Line separator, <400> field identifier
L:1126 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:52
L:1171 M:283 W: Missing Blank Line separator, <400> field identifier
L:1175 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:54
L:1240 M:283 W: Missing Blank Line separator, <400> field identifier
L:1250 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:57
M:340 Repeated in SeqNo=57
L:1281 M:283 W: Missing Blank Line separator, <400> field identifier
L:1283 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:59
M:340 Repeated in SeqNo=59
L:1326 M:283 W: Missing Blank Line separator, <400> field identifier
L:1328 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:61
L:1349 M:283 W: Missing Blank Line separator, <400> field identifier
L:1363 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:62
L:1374 M:283 W: Missing Blank Line separator, <400> field identifier
L:1388 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:63
L:1529 M:283 W: Missing Blank Line separator, <400> field identifier
L:1531 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:70
L:1576 M:283 W: Missing Blank Line separator, <400> field identifier
L:1580 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:72
L:1601 M:283 W: Missing Blank Line separator, <400> field identifier
L:1603 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:73
L:1672 M:283 W: Missing Blank Line separator, <400> field identifier
L:1695 M:283 W: Missing Blank Line separator, <400> field identifier
L:1720 M:283 W: Missing Blank Line separator, <400> field identifier
L:1777 M:283 W: Missing Blank Line separator, <400> field identifier

VERIFICATION SUMMARY
PATENT APPLICATION: US/09/606,977

DATE: 07/12/2000
TIME: 11:29:17

Input Set : D:\Maizestcs.rpt
Output Set: N:\CRF3\07122000\I606977.raw

L:1800 M:283 W: Missing Blank Line separator, <400> field identifier
L:1823 M:283 W: Missing Blank Line separator, <400> field identifier
L:1934 M:283 W: Missing Blank Line separator, <400> field identifier
L:1975 M:283 W: Missing Blank Line separator, <400> field identifier
L:2018 M:283 W: Missing Blank Line separator, <400> field identifier
L:2107 M:283 W: Missing Blank Line separator, <400> field identifier